## **RAW SEQUENCE LISTING**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number:	10/542,558
Source:	TFWP
Date Processed by STIC:	09/22/2000

## ENTERED



**IFWP** 

RAW SEQUENCE LISTING DATE: 09/22/2006
PATENT APPLICATION: US/10/542,558 TIME: 15:25:26

Input Set : A:\00530-114US1.txt \*\* \*\*

4 <110> APPLICANT: Chen, Lan Bo

Output Set: N:\CRF4\09222006\J542558.raw

```
Auclair, Daniel
 5
         Gong, Yuhong
         Dai, Meiru
 9 <120> TITLE OF INVENTION: BCRM-1 GENES AND USES THEREOF
12 <130> FILE REFERENCE: 00530-114US1
14 <140 > CURRENT APPLICATION NUMBER: US 10/542,558
15 <141> CURRENT FILING DATE: 2005-07-18
18 <150> PRIOR APPLICATION NUMBER: PCT/US2004/005338
10 1151> PRIOR FILING DATE: 2004-02-24
21 <150> PRIOR APPLICATION NUMBER: US 60/450,067
22 <151> PRIOR FILING DATE: 2003-02-25
24 <160> NUMBER OF SEQ ID NOS: 9
26 <170> SOFTWARE: FastSEQ for Windows Version 4.0
28 <210> SEQ ID NO: 1
29 <211> LENGTH: 1273
30 <212> TYPE: DNA
31 <213> ORGANISM: Homo sapiens
33 <400> SEQUENCE: 1
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35 teteetttaa gatacaggaa gettggaage ggagtettgt aagttteage eeetteetaa
                                                                          120
36 tttttagtgt cctataataa caaaaccaga catccagcta accttgcatc tctccttttg
                                                                         180
37 aaqqcaacaq tgcatcccga cagcagcaac ctgatcccca agctttttcg acctgcagcg
                                                                         240
38 ttcctqcctt tcatggcgcc cacggtattt ttgtcaatga cgccactgaa agggatcaag
                                                                          300
39 teogtgattt tacctcaggt tttcctctgt gectacatgg cagegtteaa cageatcaat
                                                                          360
                                                                         420
40 ggaaacagaa gttacacttg taagccacta gaaagatcat tactaatggc gggagccgtt
41 qcttcttcaa ctttcttagg agtaatccct cagtttgtcc agatgaagta tggcctgact
                                                                         480
42 ggcccttgga ttaaaagact cttacctgtg atcttcctcg tgcaagccag tggaatgaat
                                                                          540
43 qtctacatgt cccgaagtct tgaatccatt aaggggattg cggtcatgga caaggaaggc
                                                                          600
44 aatqtcctqq gtcattccaq aattgctqqq acaaaggctq ttagagaaac gctagcatcc
                                                                          660
45 agaatagtgc tgtttgggac ctcagctctg attcctgaag tcttcaccta cttttttaaa
                                                                          720
46 aggacccagt atttcaggaa aaacccaggg tcattgtgga ttttgaaact gtcttgtact
                                                                          780
47 gtcctggcaa tgggactgat ggtgccattt tcttttagta tatttccaca gattggacag
                                                                          840
48 atacagtact gtagtcttga agagaaaatt cagtctccaa cagaagaaac agaaatcttt
                                                                          900
49 tatcacagag gggtgtaggc gtgagtttta ggtgaattta tgtggttcct gcttgaaaac
                                                                          960
50 cttcccctct ccaggttcgg tttagagaac tttgccacag gtcttctggg gaccccagag
                                                                         1020
51 gtgtctgtgc tgacaaggcg acttcagatt ccatactgag atcgttccca ggctggcgtc
                                                                         1080
52 totggggttt ttaaggotgg otggagaaga cagtgggagg gtgccccgtc tgacacccct
                                                                         1140
53 ggggttgctg agggaacggt tggagtgggg atcggcctgc gaaaggatac tgtgaaatca
                                                                         1200
54 ctaattaact aataaacctg tctcaagttg aggatttaag ggaggtcaaa ctttttttt
                                                                         1260
                                                                         1273
55 ttttttttt ttt
57 <210> SEO ID NO: 2
58 <211> LENGTH: 221
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59 <212> TYPE: PRT
60 <213> ORGANISM: Homo sapiens
62 <400> SEQUENCE: 2
63 Met Ala Pro Thr Val Phe Leu Ser Met Thr Pro Leu Lys Gly Ile Lys
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65 Ser Val Ile Leu Pro Gln Val Phe Leu Cys Ala Tyr Met Ala Ala Phe
              20
                                   25
67 Asn Ser Ile Asn Gly Asn Arg Ser Tyr Thr Cys Lys Pro Leu Glu Arg
                               40
69 Ser Leu Leu Met Ala Gly Ala Val Ala Ser Ser Thr Phe Leu Gly Val
71 Ile Pro Gln Phe Val Gln Met Lys Tyr Gly Leu Thr Gly Pro Trp Ile
                       70
                                           75
73 Lys Arg Leu Leu Pro Val Ile Phe Leu Val Gln Ala Ser Gly Met Asn
                                       90
75 Val Tyr Met Ser Arg Ser Leu Glu Ser Ile Lys Gly Ile Ala Val Met
77 Asp Lys Glu Gly Asn Val Leu Gly His Ser Arg Ile Ala Gly Thr Lys
                               120
                                     125
          115
79 Ala Val Arg Glu Thr Leu Ala Ser Arg Ile Val Leu Phe Gly Thr Ser
                           135
81 Ala Leu Ile Pro Glu Val Phe Thr Tyr Phe Phe Lys Arg Thr Gln Tyr
                       150
83 Phe Arg Lys Asn Pro Gly Ser Leu Trp Ile Leu Lys Leu Ser Cys Thr
                                       170
                   165
85 Val Leu Ala Met Gly Leu Met Val Pro Phe Ser Phe Ser Ile Phe Pro
              180
                                   185
87 Gln Ile Gly Gln Ile Gln Tyr Cys Ser Leu Glu Glu Lys Ile Gln Ser
          195
                               200
89 Pro Thr Glu Glu Thr Glu Ile Phe Tyr His Arg Gly Val
                           215
92 <210> SEQ ID NO: 3
93 <211> LENGTH: 15
94 <212> TYPE: PRT
95 <213> ORGANISM: Artificial Sequence
97 <220> FEATURE:
98 <223> OTHER INFORMATION: Synthetically generated peptide
100 <400> SEQUENCE: 3
101 Gly Lys Arg Thr Gln Tyr Phe Arg Lys Asn Pro Gly Ser Leu Trp
102 1
104 <210> SEQ ID NO: 4
105 <211> LENGTH: 23
106 <212> TYPE: DNA
107 <213> ORGANISM: Artificial Sequence
109 <220> FEATURE:
110 <223> OTHER INFORMATION: Primer
112 <400> SEQUENCE: 4
113 acctggagag gggaaggttt tca
115 <210> SEQ ID NO: 5
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Input Set : A:\00530-114US1.txt

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116	<211> LENGTH: 19	
117	<212> TYPE: DNA	
118	<213> ORGANISM: Artificial Sequence	
120	<220> FEATURE:	
121	<223> OTHER INFORMATION: Primer	
	<400> SEQUENCE: 5	
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126	<210> SEQ ID NO: 6	
127	<211> LENGTH: 22	
128	<212> TYPE: DNA	
129	<213> ORGANISM: Artificial Sequence	
131	<220> FEATURE:	
132	<223> OTHER INFORMATION: Primer	
134	<400> SEQUENCE: 6	
135	ggtgaaggtc ggagtcaacg gt	22
137	<210> SEQ ID NO: 7	
138	<211> LENGTH: 24	
139	<212> TYPE: DNA	
140	<pre>&lt;213   ORGANICM: Aptificial Sequence</pre>	wedge East
142	<220> FEATURE:	* ***
143	<223> OTHER INFORMATION: Primer	
145	<400> SEQUENCE: 7	
146	ctggaagatg gtgatgggat ttcc	24
148	<210> SEQ ID NO: 8	
149	<211> LENGTH: 39	
150	<212> TYPE: DNA	
151	<213> ORGANISM: Artificial Sequence	
153	<220> FEATURE:	
154	<223> OTHER INFORMATION: Primer	
156	<400> SEQUENCE: 8	
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159	<210> SEQ ID NO: 9	
160	<211> LENGTH: 30	
161	<212> TYPE: DNA	
162	<213> ORGANISM: Artificial Sequence	
164	<220> FEATURE:	
165	<223> OTHER INFORMATION: Primer	
167	<400> SEQUENCE: 9	
168	gggtcgacct acacccctct gtgataaaag	30

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VERIFICATION SUMMARY

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